

FIG.1

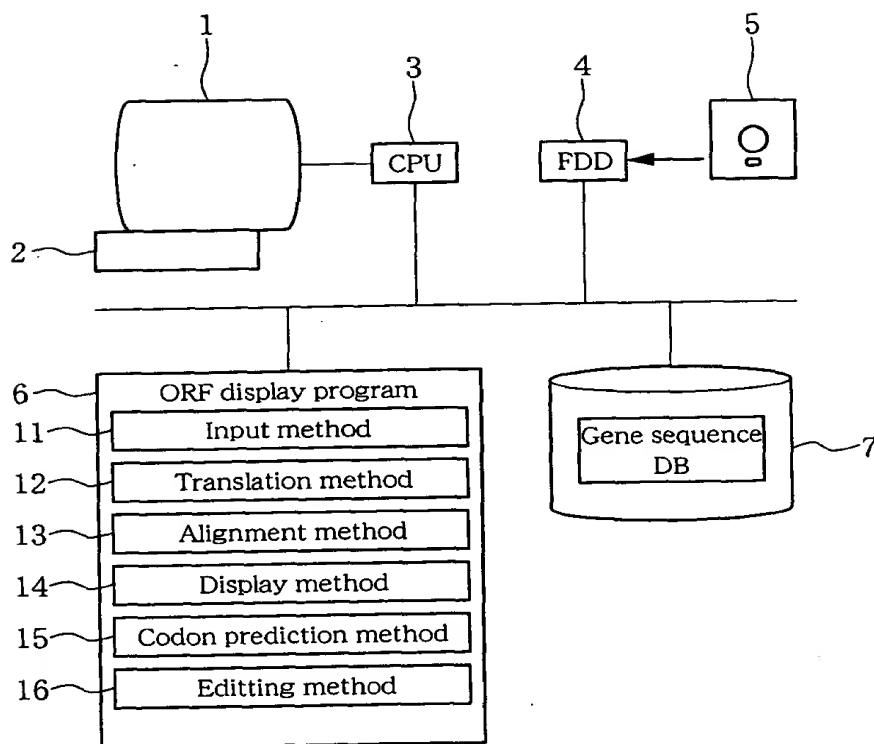


FIG.2

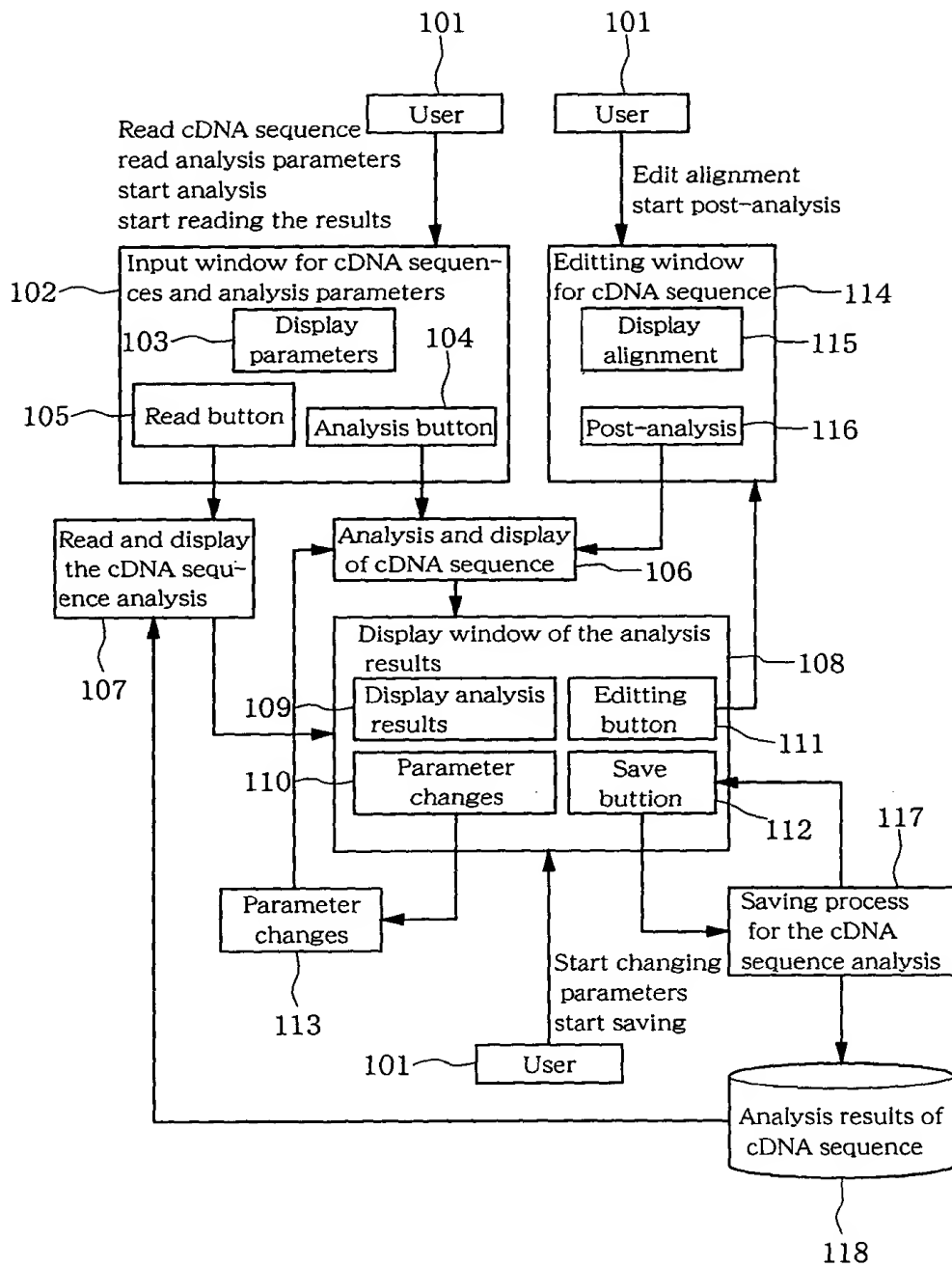


FIG.3

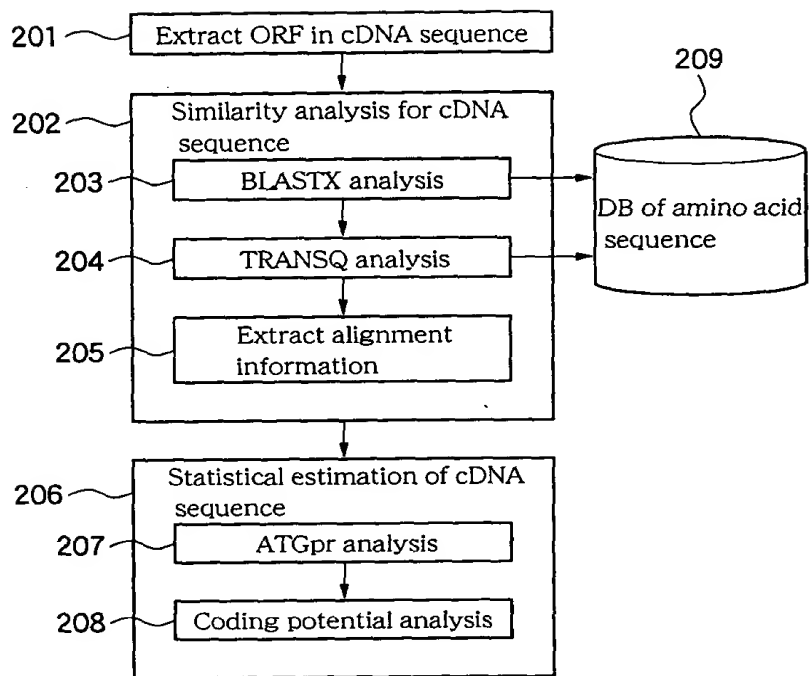


FIG.4

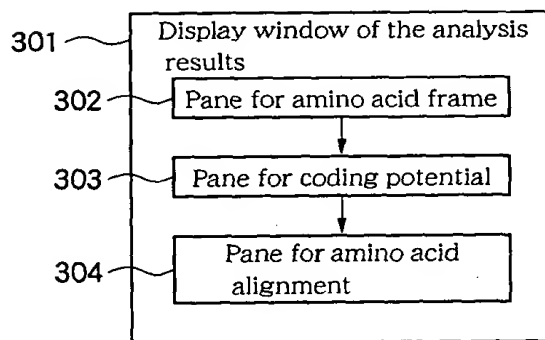


FIG.5

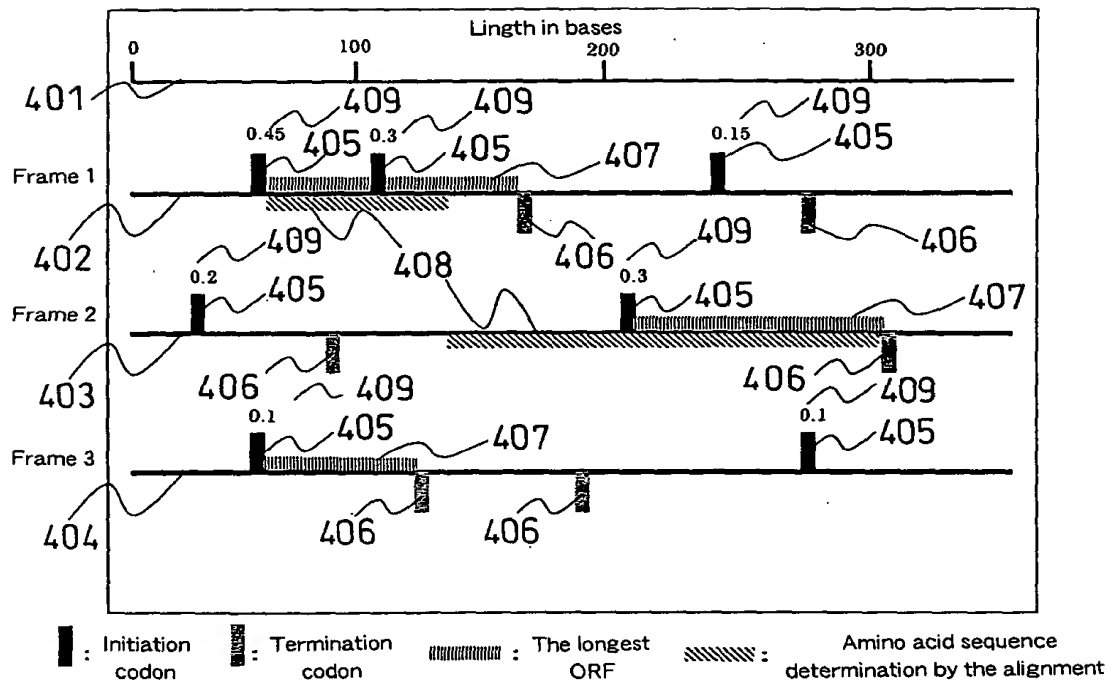


FIG.6

T06280"49304660

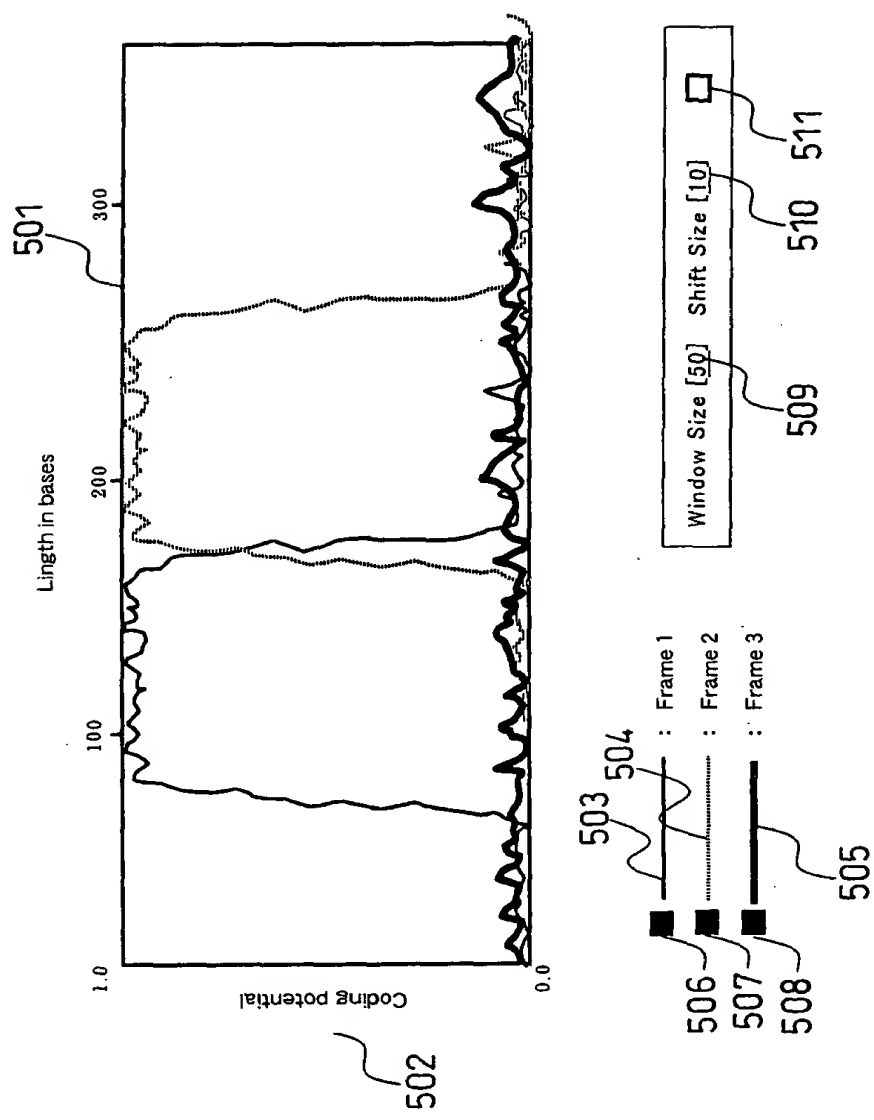


FIG.7

0940664-082901

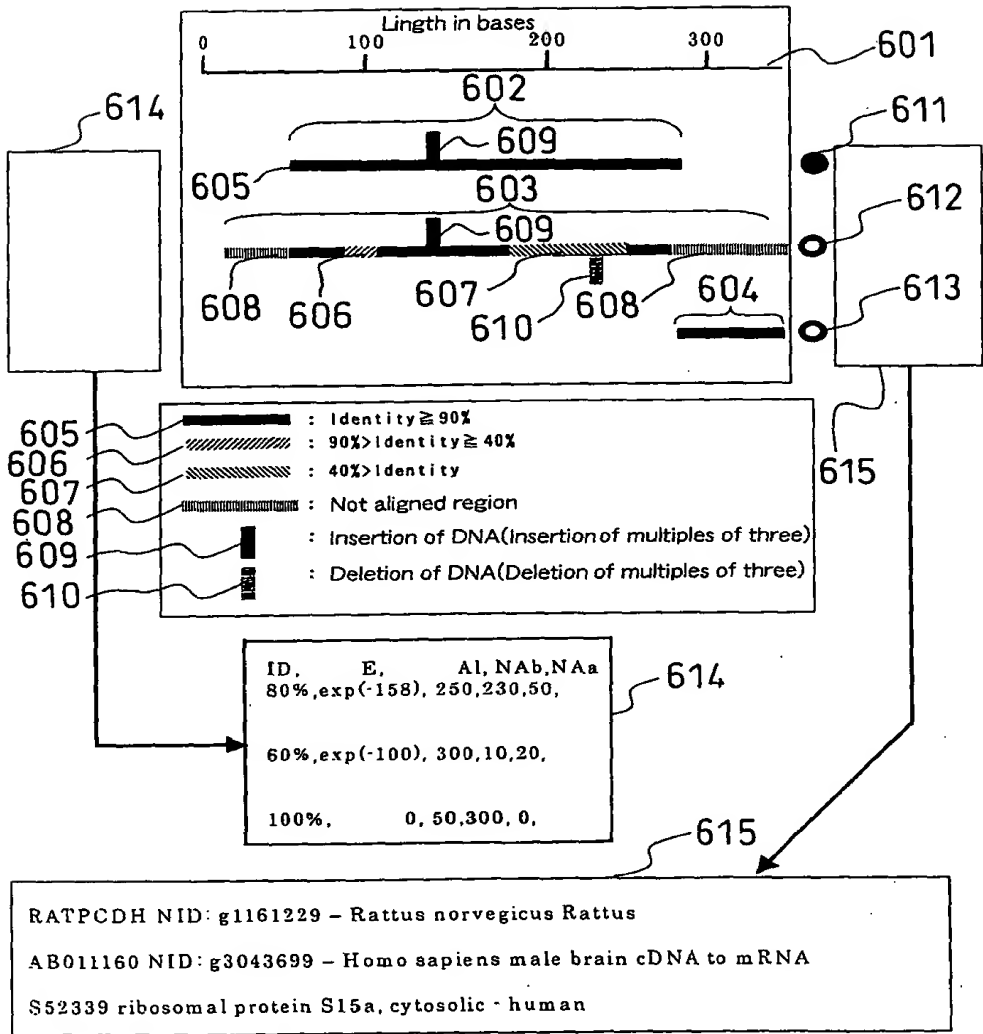


FIG.8

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Query sequence:>C-HEMBA1001174
 Target sequence:>|cl|ARL5_RAT ADP-RIBOSYLATION FACTOR-LIKE PROTEIN
 5. - RATTUS NORVEGICUS (RAT)
 Score: 794 + strand
 Alignment region Query : 208 .. 744
 Target: 1 .. 179
 Identity: 144/179(80%) Positive: 165/179(92%)

60 :	70 :	80 :	90 :	100 :	110
Query: 208 ATGgggctgatcttcgccaaactgtggagcctcttctgtaaccaagaacacaaaagtaatt					
MetGlyLeuIlePheAlaLysLeuTrpSerLeuPheCysAsnGlnGluHisLysValIle					
: : . : : :					
Target: 1 MetGlyIleLeuPheThrArgIleTrpArgLeuPheAsnHisGlnGluHisLysValIle					

702

120 :	130 :	140 :	150 :	160 :	170
Query: 267 atagtgggactgagataatgcagggaaaaccaccattctttaccaattcttaATGaatgaa					
IleValGlyLeu AspAsnAlaGlyLysThrThrIleLeuTyrGlnPheLeuMetAsnGlu					
Target: 20 IleValGlyLeu AspAsnAlaGlyLysThrThrIleLeuTyrGlnPheSerMetAsnGlu					

Submit Close Refresh

703

704

705

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FIG.9

